

INFORMAL SEQUENCE LISTING

NOVEL PROTEINS

PN7098

SEQ ID NO:1 >PN7098 (NT)

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PN7771

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SEQ ID NO:5 >PN9109 (NT)

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agatgtgaaatcagcaacttccctaccatcgaattttatcatgctcttggaaactca

acaaaacagctgtAACATGCTTAAGTAACAATTAGCAGTGATTCCAAAGTCGTGTTCTCAATCCTAAAACACATCACCCCTACCTGAGTTAACCTAGCACACCAACTTGTACTTTCCTCTTGGTGTTCAGTAAGCAAAGAAGATCCAATTTCCTCTTTCAAACATCATGAGTATTAGCTTAGATAAAAGTTCTTAT

SEQ ID NO:10 >PN13734 (AA)

MDRILPDGGVAKTMMESSLADFMQEVGYGFCA\$IECRNIIIVQFGVREVTAAQVARVLGMMARTHSGLT\$DGIPLO\$ISAPGSGIWSDGKDKSDGAQAHTWNVEVLIDVLKELNPSLNFKEVTYELDHPGFQIRDSKGLHNVVYGIQRGLGM\$EVFPVDLIYRPWKHAEGQLSFIQHSLINPEIFCFADYPCHTVATDILKAPPEDDNREIA\$TWKSLDLIESLLRLAEVGQYEQVKQLFSFPIKHCPDMVLALLQINTSWHTLRHELISTLMP\$IFLGNHPNSAIILHYAWHGQGQSPSIRQLIMHAMAEWYMRGEQYDQAKLSRILDVAQDLKALSM\$LLNGTPFAFVIDLAALASREYLKLDKWLT\$DKIREHGE\$PIQACMTFLKRRCP\$ILGG\$LAPEKDQPKSAQLP\$PETLATMLACLQACAGSVSQELSETILTMVANC\$NVMNKARQPPGVMPKGRPPSASSLDAISPVQIDPLAGMTSLSIGGSAA\$PHTQSMQGFP\$NLGSAFSTPQSPA\$KA\$FPP\$LP\$TNQ\$PTFQQSKMKPSDLSQVWPELPVGG\$LGTS\$LTGIGT\$GALGLPAVNNDP\$FVQRKLGT\$GLNQ\$PTFQQSKMKPSDLSQVWPEANQHFSKEIDDEANSYFQRIYNHPPHPTMSVDEVLEMLQRFKD\$TIKREREVFNCMLRNLFEEYRFFPQY\$PD\$KELHI\$ACLF\$GGIEKGLV\$TYMALGLALRYVLEALRK\$PFGSKMYYFGIAALDRFKNRLKDYPQYCQHLASI\$HFMQFPHLQEYIEYGQ\$SRDPPV\$KM\$QGSITTPGSIALAQQAQAQVPAKAPLAGQVSTMVTTSTTVAKTV\$TRPTGV\$F\$KKDVPPSINTTNDTLLVATDQTERI\$VEP\$ENI\$QEKIAFIFNNL\$SQSNMTQ\$KVEEL\$KETVKEEFMPWV\$QYLVMKRV\$IEPNFH\$LYSNFLDTLK\$NPEFN\$KMVLNEY\$YNIKVLLTSDKAAANFSDRSLLKNGHWLGMITLAKNK\$PILH\$TDLV\$KSLL\$EAYVKGQ\$ELLYVVPFVAKVLESSI\$RSVVFRPPNPWTMAIMNVLAELHQEHDLKLN\$KFEIEVLCKNL\$ALDINELKPGNLL\$KD\$K\$DL\$K\$NL\$DEQLSAPKKDV\$KQPEELPPITT\$TT\$PATNTTCTATVPPQ\$QYSYHDINVYSLAGLAPHITLNPTIPLFQAH\$PQLKQCVRQAIERAVQELVHPV\$DRSIKIAMTCEQIVRKDFALDSEESRMRIAHHMMRNLTAGMAMITCREPLLMSISTNL\$KNSFASALRTASPQ\$QREMMDQAAQLAQDNCELACCFIQKTAVEKAGPEMDKRLATEFELRKHARQEGR\$RYCDPV\$VLT\$YQAERMPEQIRLK\$VGG\$VDP\$KQLAVYEEFARNVPGFLPTNDL\$QPTGFLAQPMKQAWATDDVAQIYDKCITELEQHLHAI\$PPTLAMNPQ\$AQALRS\$LLEV\$V\$VLSRN\$RDIAALGLL\$Q\$AVEGL\$DATSGADADLLLRYRECHLLVLKALQDGRAY\$GSPWCNKQITRC\$IECRDEY\$KYNVEAVELLIRNH\$LVNMQQYDLH\$LAQSMENGLNYMAFA\$AMQLV\$KILLV\$DERSVAHVTEADLFHTIE\$TL\$MRINAHSGNAPEGLPQLMEVVR\$NSYEAMIDRAHGG\$PNFMMHSGI\$SQASEYDDPPGLREKA\$EYL\$LREWVNLYHSAAGRDSTKAFSAFVGQ\$MHQQGILKTD\$DLITRF\$FRLCTEMC\$VEISYRAQAEQQHNPAANPTMIRAKCYHNLD\$AFVRLIALLVKHS\$GEATNTVTKINLLNKVLGIVVGVLLQDH\$DVRQ\$SEFQQLPYHRIFIMLL\$E\$LN\$APEHVLETINFQ\$TLAFCNTFHILRPTKAPGFVYAWLELI\$SHRIFI\$IA\$MLAHTPQQKGWP\$MYAQLLIDL\$F\$KYLAPFLRN\$VELTKPMQILYKGT\$LRVLLVLLHDFPEFLCDYHYGFCDVIPPNCIQLRNLL\$IAFPRNMRLPDPFTPNLKV\$DMLSEINIAPRILTNFTGV\$MPPQF\$KKD\$LSY\$LKTRSPVTF\$LS\$DL\$RSNLQVSNEPGNRYNLQLINALVLYVGTQAI\$IAH\$HNGSTPSMSTI\$THSAHMDIFQNLAVDLDTEGRYLF\$LN\$IA\$ANQLRYPNSHTHYFSCTMLYLF\$AEANTEAIQEQITRVLLERLIVNRPHPWGLLITFIELIKNPAFK\$FWNHEFVHCAPEIEKLFQ\$VAQCCMGQKQAQQVMEGT\$GAS

Antisense Oligonucleotides Directed Against:

ERK3

SEQ ID NO:11 5'-ttctctgccatTTgaaacc-3'

SEQ ID NO:12 5'-ctctgccatTTgaaaccct-3'

SEQ ID NO:13 5'-ctgccatTTgaaacccta-3'

PRKAR1A

SEQ ID NO:14 5'-ccagactccatggTCTCTG-3'

SEQ ID NO:15 5'-agactccatggTCTCTGGG-3'

SEQ ID NO:16 5'-actccatggTCTCTGGGA-3'

KRT23 (209)

SEQ ID NO:17 5'-tgcttctccatttcttgggg-3'

SEQ ID NO:18 5'-cttctccatttcttgggg-3'

SEQ ID NO:19 5'-tctccatttcttgggg-3'

PN7098

SEQ ID NO:20 5'-ttAGCCACCATGGAGAGCAA-3'

SEQ ID NO:21 5'-AGCCACCATGGAGAGCAATT-3'

SEQ ID NO:22 5'-CCACCATGGAGAGCAATTAG-3'

AL117237

SEQ ID NO:23 5'-ttCCTCAGCATAGATTTAT-3'

SEQ ID NO:24 5'-CCTCAGCATAGATTTATGA-3'

SEQ ID NO:25 5'-TCAGCATAGATTTATGAGA-3'

PCNT2

SEQ ID NO:26 5'-tcaactccatctccctcag-3'

SEQ ID NO:27 5'-aactccatctccctcagac-3'

SEQ ID NO:28 5'-cttccatctccctcagactc-3'

PROX1

SEQ ID NO:29 5'-tggtcaggcatcactggacg-3'

SEQ ID NO:30 5'-gtcaggcatcactggacggt-3'

SEQ ID NO:31 5'-caggcatcactggacggta-3'

HOOK1

SEQ ID NO:32 5'-gtctcctccatggccggcgc-3'

SEQ ID NO:33 5'-ctcctccatggccggcgccg-3'

SEQ ID NO:34 5'-cctccatggccggcgccgcc-3'

IGHG1

No ATG identified. Used three different sequences, one from the beginning, middle, and end.

SEQ ID NO:35 5'-gaggagggtgccaggggaa-3'

SEQ ID NO:36 5'-cagggtcttcgtggctacg-3' :

SEQ ID NO:37 5'-ctgcgtgttagtggttgtca-3'

GOLGA2

SEQ ID NO:38 5'-accgactccattaccagccc-3'

SEQ ID NO:39 5'-cgactccattaccagcccct-3'

SEQ ID NO:40 5'-actccattaccagcccctga-3'

KIAA0555

SEQ ID NO:41 5'-ttcttggacattgttccttt-3'

SEQ ID NO:42 5'-cttggacattgttcctttct-3'

SEQ ID NO:43 5'-tggacattgttcctttctaa-3' :

LRPPRC

SEQ ID NO:44 5'-gcttcctccatTTTgcag-3'

SEQ ID NO:45 5'-ttcctccatTTTgccagga-3'

SEQ ID NO:46 5'-cctccatTTTgccaggaaa-3'

SET

SEQ ID NO:47 5'-tgcgccgacatggtgctgct-3'

SEQ ID NO:48 5'-cgccgacatggtgctgctcc-3'

SEQ ID NO:49 5'-ccgacatggtgctgctccgc-3'

TL21

No ATG identified. Used three different sequences, one from the beginning, middle, and end.

SEQ ID NO:50 5'-cttttcagttccttggtc-3'

SEQ ID NO:51 5'-cattttctcaagtcccttc-3'

SEQ ID NO:52 5'-taatccaggataactcagca-3'

KAISO

SEQ ID NO:53 5'-ctactctccatgcctgcttc-3'

SEQ ID NO:54 5'-actctccatgcctgcttccc-3'

SEQ ID NO:55 5'-tctccatgcctgctccgg-3'

PN7771

SEQ ID NO:56 5'-acctcagccatctgcaacaa-3'

SEQ ID NO:57 5'-ctcagccatctgcaacaagg-3'

SEQ ID NO:58 5'-cagccatctgcaacaaggct-3'

TSP3

SEQ ID NO:59 5'-tgcgtctccatgcctctcag-3'

SEQ ID NO:60 5'-cgtctccatgcctctcagcc-3'

SEQ ID NO:61 5'-tctccatgcctctcagccgg-3'

MDH1

No ATG identified. Used three different sequences, one from the beginning, middle, and end.

SEQ ID NO:62 actctgattggttcagacat

SEQ ID NO:63 5'-cagtcacaccaagttaaga-3'

SEQ ID NO:64 5'-ttgattacaacagggaatga-3'

GA17

SEQ ID NO:65 5'-gggacgctcatggtgcgaggc-3'

SEQ ID NO:66 5'-gacgctcatggtgcgaggccc-3'

SEQ ID NO:67 5'-cgctcatggtgcgaggccgg-3'

CAPN4

SEQ ID NO:68 5'-accaggaacatggctgac-3'

SEQ ID NO:69 5'-caggaacatggctgactc-3'

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SEQ ID NO:70 5' -ggaacatggctgcgactcac-3'

BAT3

SEQ ID NO:71 5' -ttaggctccatggccgacag-3'

SEQ ID NO:72 5' -aggctccatggccgacaggt-3'

SEQ ID NO:73 5' -gctccatggccgacaggtct-3'

ABLIM1

SEQ ID NO:74 5' -aaggcaggcatttggcatg-3'

SEQ ID NO:75 5' -ggcaggcatttggcatgga-3'

SEQ ID NO:76 5' -caggcatttggcatggatt-3'

NICE-4

SEQ ID NO:77 5' -gatgtcatcattacagtat-3'

SEQ ID NO:78 5' -tgtcatcattacagtatat-3'

SEQ ID NO:79 5' -tcatcattacagtatac-3'

CYT4

SEQ ID NO:80 5' -cacaggtccattccgacgcc-3'

SEQ ID NO:81 5' -caggtccattccgacgcctc-3'

SEQ ID NO:82 5' -ggtccattccgacgcctctg-3'

JNK3A2

SEQ ID NO:83 5' -tggaggctataaataaccac-3'

SEQ ID NO:84 5' -gaggctataaataaccacag-3'

SEQ ID NO:85 5' -ggctataaataaccacagct-3'

C-NAP1

SEQ ID NO:86 5'-cttgtctccatgagggcacc-3'

SEQ ID NO:87 5'-tgtctccatgagggcaccag-3'

SEQ ID NO:88 5'-tctccatgagggcaccaggt-3'

VCL(1066)

SEQ ID NO:89 5'-aacactggcatcgcgccggc-3'

SEQ ID NO:90 5'-cactggcatcgcgccggcga-3'

SEQ ID NO:91 5'-ctggcatcgcgccggcgagc-3'

PSF

SEQ ID NO:92 5'-tccc gagacatgtctgtggt-3'

SEQ ID NO:93 5'-ccgagacatgtctgtggtca-3'

SEQ ID NO:94 5'-gagacatgtctgtggtcaag-3'

FNTA

There are only 6 nucleotides before the start codon, so I used 11 nucleotides after the start codon to maintain the 20 nuclueotides and I didn't shift left.

SEQ ID NO:95 5'-cctcggtggccgcacatctcg-3'

SEQ ID NO:96 5'-tcggtgccgcacatctcgcc-3'

TPRD

SEQ ID NO:97 5'-aaattgtccatggtgacaaa-3'

SEQ ID NO:98 5'-attgtccatggtgacacaagt-3'

SEQ ID NO:99 5'-tgtccatggtgacacaagtct-3'

PN9190

No ATG identified. Used three different sequences, one from the beginning, middle, and end.

SEQ ID NO:100 5'-ctataactactactgtgcac-3'

SEQ ID NO:101 5'-gtactcttcttaaccagtg-3'

SEQ ID NO:102 5'-ggctctgtgtctggggac-3'

PPL

There are only 11 nucleotides identified before the start codon so I didn't shift to the left.

SEQ ID NO:103 5'-agcgagttcatggtggcgct-3'

SEQ ID NO:104 5'-cgagttcatggtggcgctcg-3'

Golgin-84

SEQ ID NO:105 5'-aaccaagacatgatggcagc-3'

SEQ ID NO:106 5'-ccaagacatgatggcagcag-3'

SEQ ID NO:107 5'-aagacatgatggcagcagga-3'

CLIC1

SEQ ID NO:108 5'-cacagtaccatgaacacgtct-3'

SEQ ID NO:109 5'-cagtaccatgaacacagtctct-3'

SEQ ID NO:110 5'-gtaccatgaacacagtctctgg-3'

AKR7A2

SEQ ID NO:111 5'-ggccgggacatggcgagcgc-3'

SEQ ID NO:112 5'-ccgggacatggcgagcgcgc-3'

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SEQ ID NO:113 5' -gggacatggcgagcgcgcg-3'

UNR (798)

SEQ ID NO:114 5' -tcaaagctcatctcgcagtg-3'

SEQ ID NO:115 5' -aaagctcatctcgcagtgat-3'

SEQ ID NO:116 5' -agctcatctcgcagtgatac-3'

GBDR1

SEQ ID NO:117 5' -tgcacgaacatcccgccgccc-3'

SEQ ID NO:118 5' -cacgaacatcccgccgccc-3'

SEQ ID NO:119 5' -cgaacatccgcccggccg-3'

KIAA0614

No nucleotides were identified before the start codon, so I chose three different 20 nucleotide sequences from the beginning, middle, and end.

SEQ ID NO:120 5' -actgttatctccttccaccat-3'

SEQ ID NO:121 5' -gattgactttcttacaaga-3'

SEQ ID NO:122 5' -gaacatctccagggcccccc-3'

LDHM

SEQ ID NO:123 5' -agagttgccatattggactt-3'

SEQ ID NO:124 5' -agttgccatattggacttg-3'

SEQ ID NO:125 5' -ttgccatattggacttgaa-3'

EIF3S10

SEQ ID NO:126 5' -taggccccatcttggcgcc-3'

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SEQ ID NO:127 5'-ggccggcatcttggcggcag-3'

SEQ ID NO:128 5'-ccggcatcttggcggcaggc-3'

SLAP2

SEQ ID NO:129 5'-ttttttccatttcatttat-3'

SEQ ID NO:130 5'-ttttccatttcatttat-3'

SEQ ID NO:131 5'-tttccatttcatttat-3'

SART-1

SEQ ID NO:132 5'-gacgaccataggcacg-3'

SEQ ID NO:133 5'-cgaccataggcacgtc-3'

SEQ ID NO:134 5'-accataggcacgtcca-3'

ITRAF

SEQ ID NO:135 5'-tttttatccattcctttcg-3'

SEQ ID NO:136 5'-tttatccattcctttcg-3'

SEQ ID NO:137 5'-tatccattcctttcg-3'

NUMA1

SEQ ID NO:138 5'-tggagtgtcatcttggtgat-3'

SEQ ID NO:139 5'-gagtgtcatcttggtgatgc-3'

SEQ ID NO:140 5'-gtgtcatcttggtgatgcca-3'

SPA1

SEQ ID NO:141 5'-cacatggcatgctgtgg-3'

SEQ ID NO:142 5'-catggcatgctgtggc-3'

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SEQ ID NO:143 5'-tgggcatgctctgtggcct-3'

FYCO1

SEQ ID NO:144 5'-gtggaggccatggtgagttt-3'

SEQ ID NO:145 5'-ggaggccatggtgagttgc-3'

SEQ ID NO:146 5'-aggccatggtgagttgcct-3'

DNCL1

SEQ ID NO:147 5'-cggtcgcacatggttaccgt-3'

SEQ ID NO:148 5'-gtcgcacatggttaccgtgg-3'

SEQ ID NO:149 5'-cgcacatggttaccgtggag-3'

ACTB

SEQ ID NO:150 5'-tcatcatccatggtgagctg-3'

SEQ ID NO:151 5'-atcatccatggtgagctggc-3'

SEQ ID NO:152 5'-catccatggtgagctggcgg-3'

KU70

SEQ ID NO:153 5'-caccctgacatgttggctac-3'

SEQ ID NO:154 5'-ccctgacatgttggctactg-3'

SEQ ID NO:155 5'-ctgacatgttggctactgct-3'

COPP

SEQ ID NO:156 5'-cgcagaggcatggtgcgctcg-3'

SEQ ID NO:157 5'-cagaggcatggtgcgctcggt-3'

SEQ ID NO:158 5'-gaggcatggtgcgctcggtcc-3'

OSTP

SEQ ID NO:159 5' -gcaattctcatggtagttag-3'

SEQ ID NO:160 5' -aattctcatggtagttagttt-3'

SEQ ID NO:161 5' -ttctcatggtagttagtttt-3'

SLC8A1

SEQ ID NO:162 5' -atgttgtacatgacacttcc-3'

SEQ ID NO:163 5' -gttgtacatgacacttccaa-3'

SEQ ID NO:164 5' -tgtacatgacacttccaact-3'

A2-CAT

SEQ ID NO:165 5' -gccgaagtcatgctccctct-3'

SEQ ID NO:166 5' -cgaagtcatgctccctctgt-3'

SEQ ID NO:167 5' -aagtcatgctccctctgtgg-3'

THRSP

Only 10 nucleotides were identified before the start codon, so I included 7 nucleotides after the start codon and I didn't shift left two spots.

SEQ ID NO:168 5' -tagcacctgcatggttgctt-3'

SEQ ID NO:169 5' -gcacctgcatggttgcttcc-3'

OPA1

SEQ ID NO:170 5' -agtgcacatccccggggc-3'

SEQ ID NO:171 5' -tcgcacatccccggggcgg-3'

SEQ ID NO:172 5' -gccacatccccggggcgcccc-3'

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PN12218

SEQ ID NO:173 5' -gcggagcagtgacgcggcat-3'

SEQ ID NO:174 5' -gaggcctttccacagaaaa-3'

SEQ ID NO:175 5' -atgctgctgctcagctgcat-3'

CIN85

SEQ ID NO:176 5' -ggcctccaccattgcgtcga-3'

SEQ ID NO:177 5' -atggcctccaccattgcgtc-3'

SEQ ID NO:178 5' -ctatggcctccaccattgcg-3'

PN13734

SEQ ID NO:179 5' -gatcctgtccattagaatgt-3'

SEQ ID NO:180 5' -aggatcctgtccattagaat-3'

SEQ ID NO:181 5' -gcaggatcctgtccattaga-3'

FUBP1

SEQ ID NO:182 5' -atagtctgccatggttgcac-3'

SEQ ID NO:183 5' -gaatagtctgccatggttgc-3'

SEQ ID NO:184 5' -ttgaatagtctgccatggtt-3'

LRP1

SEQ ID NO:185 5' -cggggtcagcatggtgtgg-3'

SEQ ID NO:186 5' -ggcggttcagcatggtgtg-3'

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SEQ ID NO:187 5' -acggcggttcagcatggtg-3'

FUS(526) and FUS(525)

SEQ ID NO:188 5' -gttgaggccatgtccgcgc-3'

SEQ ID NO:189 5' -tcgttgaggccatgtccgc-3'

SEQ ID NO:190 5' -aatcgttgaggccatgtcc-3'

XE169

SEQ ID NO:191 5' -ccccggctccatggtgggcc-3'

SEQ ID NO:192 5' -gaccggctccatggtggg-3'

SEQ ID NO:193 5' -cggaccggctccatggtg-3'

NAG4

SEQ ID NO:194 5' -cttcttgcctatccgacc-3'

SEQ ID NO:195 5' -tgcttcttgcctatccga-3'

SEQ ID NO:196 5' -tgtgcttgcctatcc-3'

ESRRA

SEQ ID NO:197 5' -gagctcatctccaatcccatt-3'

SEQ ID NO:198 5' -agcagggaaaggggccggga-3'

SEQ ID NO:199 5' -gggccagcacttgcggct-3'

CTNNB1

SEQ ID NO:200 5' -tttagtagccattgtccacg-3'

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SEQ ID NO:201 5' -gcttgagtagccattgtcca-3'

SEQ ID NO:202 5' -cagcttgagtagccattgtc-3'

CIT

SEQ ID NO:203 5' -gaacttcaacatctccccac-3'

SEQ ID NO:204 5' -ttgaacttcaacatctcccc-3'

SEQ ID NO:205 5' -atttgaacttcaacatctcc-3'

HSPB1

SEQ ID NO:206 5' -aaagaggctattccgtac-3'

SEQ ID NO:207 5' -tcaaagaggctattccgt-3'

SEQ ID NO:208 5' -gatcaaagaggctattccc-3'

ROCK2

SEQ ID NO:209 5' -gggcggctcatgccgccac-3'

SEQ ID NO:210 5' -ggggccggctcatgccgcc-3'

SEQ ID NO:211 5' -gcggggccggctcatgccg-3'

TILP (392)

SEQ ID NO:212 5' -ctaccagtgcattccatgaca-3'

SEQ ID NO:213 5' -ttctaccagtgcattcatga-3'

SEQ ID NO:214 5' -tcttctaccagtgcattccat-3'

LRRFIP2a

SEQ ID NO:215 5' -aggagtccccatcttgtgtt-3'

SEQ ID NO:216 5' -gcaggagtccccatcttgtg-3'

SEQ ID NO:217 5' -aagcaggagtccccatcttg-3'

PSAP

SEQ ID NO:218 5' -gagggcgtacatagcaccgt-3'

SEQ ID NO:219 5' -aagagggcgtacatagcacc-3'

SEQ ID NO:220 5' -ggaagagggcgtacatagca-3'

NOTCH2

SEQ ID NO:221 5' -cagggcgcccattttctcg-3'

SEQ ID NO:222 5' -cgcagggcgcccattttctc-3'

SEQ ID NO:223 5' -ggcgcagggcgcccatttc-3'

Peptide Fragments

of PRAK Interactors

ERK3

SEQ ID NO:224 PMDEPISSHPFHIEDEVDDI
SEQ ID NO:225 TEPCWQYSDHHENKYCDLEC
SEQ ID NO:226 LISKSVSQEKGEMANLAQ
SEQ ID NO:227 GEFLFNKQLESIGIPQFHSP
SEQ ID NO:228 VWRESEVNHYEPKLIIDL
SEQ ID NO:229
QFSEHDWPVHNFDIDEVQLDPRALSDVTDEEVQVDPRKYLDGDREKYL
SEQ ID NO:230
NSGEFLFNKQLESIGIPQFHSPVGSPPLKSIQATLTPSAMKSSPQIPHQTY

PRKAR1A

SEQ ID NO:231 MESGSTAASEEARSLRECEL
SEQ ID NO:232 YVQKHNIQALLKDSIVQLCT
SEQ ID NO:233 ARPERPMAFLREYFERLEKE
SEQ ID NO:234 EAKQIQNLQKAGTRTDSRED
SEQ ID NO:235 EISPPPPNPVVKGRRRGAI
SEQ ID NO:236
EAKQIQNLQKAGTRTDSREDEISPPPPNPVVKGRRRGAI SAEVYTEEDA
SEQ ID NO:237
AASEEARSLRECELYVQKHNIQALLKDSIVQLCTARPERPMAFLREYFER

KRT29 (209)

SEQ ID NO:238 MEKHHVPSDFNVNVKVDTGP
SEQ ID NO:239 REDLIKVLEDMRQEYELIIK

PN7098

SEQ ID NO:240 DEEDYTEPVADNETDYVEVM
SEQ ID NO:241 MQAYDHLSYETPYETPQDEG
SEQ ID NO:242 EITKPKRIRPSFKEAALRAY
SEQ ID NO:243 VCGGAGGGLYGIDSMPDLRR
SEQ ID NO:244 FKKTQLALIYPMSSTIPHNF
SEQ ID NO:245
LYGIDSMPDLRRKKTLPIVRDVAMTLAARKSGLSLAMVIRTSLNNEELKM
SEQ ID NO:246
LSYETPYETPQDEGYDGPADDMVSEEGLEPLNETSAEMEIREDENQNIPE

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SEQ ID NO:247 YLGLALDVDRRIKKDQEED
SEQ ID NO:248 DQGPPCPRLSRELLEVVEPE
SEQ ID NO:249 LQDSLDRCYSTPSSCLEQPD
SEQ ID NO:250 SCQPYGSSFYALEEKHVGFS

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SEQ ID NO:251
YLGLALDVDRRIKKDQEEEEDQGPPCPRLSRELLEVVEPEVLQDSLDRCYS
SEQ ID NO:252
RLSRELLEVVEPEVLQDSLDRCYSTPSSCLEQPDSCQPYGSSFYALEEKH

PCNT2

SEQ ID NO:253 TPEQRGIFTISDHPAEQRGM
SEQ ID NO:254 AVHGLELEALRLSLSNMHTA
SEQ ID NO:255 HELELLREQHAREKEEVVLR
SEQ ID NO:256 EDLQSEHGRCLLEDLEFKFKE
SEQ ID NO:257 LSELHEQOLLARTSRVEDLEQ
SEQ ID NO:258
ELEALRLSLSNMHTAQLELTQANLQKEKETALTELREMLNSRRAQELALL
SEQ ID NO:259
KELSAKHQSEMEDLQNQFQKELAEQRAELEKIFQDKNQAERALRNLESHH

PROX1

SEQ ID NO:260 QQLEDMQKQLLHVQEKFYQI
SEQ ID NO:261 ELDPGQFIDRARALIREQEM
SEQ ID NO:262 SQVVDTVVVKVFSAKPSRQVP
SEQ ID NO:263 PLQIPQARFAVNNGENHNHFHT
SEQ ID NO:264 TFGNVQMASSTDQTEALPLV
SEQ ID NO:265
FSAKPSRQVPQVFPLQIPQARFAVNNGENHNFTANQRLQCFGDVIIPNP
SEQ ID NO:266
RESYRENKRKQKLPOQQQQSFQQLVSARKEQKREERRQLKQQLEDMQKQL

HOOK1

SEQ ID NO:267 MEETQPPPQPKLPLCDSLMI
SEQ ID NO:268 WLQTFNTASPCQDVKQLTSG
SEQ ID NO:269 EELQEALAEKEELRQRCEEL
SEQ ID NO:270 IDVLRATSDKANKLESTVEI
SEQ ID NO:271 YRQKLQDLNDLRKQVKTLQE
SEQ ID NO:272
EDVGDNWRIKASNKKVLQGIMSYHEFLGQQISEALIPDLNQITECSDP
SEQ ID NO:273
RALEELQEALAEKEELRQRCEELDMQVTTLQDEKNSLVSENEMMNEKLDQ

IGHG1

SEQ ID NO:274 SVFLFPPPKDTLMISRTP
SEQ ID NO:275 VTCVVVDVSHEDPEVKFNWY
SEQ ID NO:276 VDGVEVHNNAKTKPREEQYNS
SEQ ID NO:277 CLVKGFYPSDIAVEWESNGQ
SEQ ID NO:278 PENNYKTTPPVLDSDGSFFL
SEQ ID NO:279
TYRVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQV

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SEQ ID NO:280

DIAVEWESNGQPENNYKTPVLDSDGSFFLYSKLTVDKSRWQQGNVFSC

GOLGA2

SEQ ID NO:281 EQQLQAEAEHLRKELEGLAG

SEQ ID NO:282 QLQAQVQDNEGLSRLNREQE

SEQ ID NO:283 QAQVQDNEGLSRLNREQEER

SEQ ID NO:284

EQQLQAEAEHLRKELEGLAGQLQAQVQDNEGLSRLNREQEERLLELERA

SEQ ID NO:285

AEHLRKELEGLAGQLQAQVQDNEGLSRLNREQEERLLELERAELWGEQA

KIAA0555

SEQ ID NO:286 AEESELRFQRLTKEYQALQR

SEQ ID NO:287 AYALLQEQTGGIIDAEREAK

SEQ ID NO:288 AQEQLQAEVLRYKAKIEDLE

SEQ ID NO:289 ATLAQKGQIEKQEAENHRLQ

SEQ ID NO:290 QELQDARDQNELLFRNLEL

SEQ ID NO:291

AEESELRFQRLTKEYQALQRAYALLQEQTGGIIDAEREAKAQEQLQAEVL

SEQ ID NO:292

RYKAKIEDLEATLAQKGQIEKQEAENHRLQQELQDARDQNELLFRNLEL

LRPPRC

SEQ ID NO:293 MEEANIQPNSRTYQRLIASY

SEQ ID NO:294 MENAENILTVMRDAGIEPGP

SEQ ID NO:295 QTLEKVEKFELHLMRDRLQ

SEQ ID NO:296 VFGSFFLQHCVTMNTPVEKL

SEQ ID NO:297 TDYCKKLKEVQMHSFPLQFT

SEQ ID NO:298

MENAENILTVMRDAGIEPGPDTYLALLNAYAEKGIDIDHVQQTLEKFE

SEQ ID NO:299

IIFSFSKAGYLSMSQKFWKKFTCERRYIPDAMNLILLLVTEKLEDVALQI

Of MAPKAP-K2 Interactors

SET(277)

SEQ ID NO:300 LTRVEVTEFEDIKSGYRIDF

SEQ ID NO:301 YFDENPYFENKVLSKEFHLN

SEQ ID NO:302 TKRSSQTQNKA SRKRQHEEP

SEQ ID NO:303 ESFFT WFTDHSDAGADELGE

SEQ ID NO:304 VIKDDIWP NPLQYYLVPMD

SEQ ID NO:305

LTRVEVTEFEDIKSGYRIDFYFDENPYFENKVLSKEFHLN ESGDPSSKST

SEQ ID NO:306

EIKWKSGKD LTKRSSQTQNKA SRKRQHEEPESFFT WFTDHSDAGADELGE

OF MAPKAP-K3 Interactors

KAISO

SEQ ID NO:307 GVGSKHLMEGQKIIITLDTAT
SEQ ID NO:308 EIEGLSTGCKVYANIGEDTY
SEQ ID NO:309 DIVIPVKDDPDEGEARLENE
SEQ ID NO:310 HEIHHTGERRYQCLACGKSF
SEQ ID NO:311 INYQFMSSHISVHSQDPSG
SEQ ID NO:312 HDDHYELIVDGRVYYICIVCKRSYVCLTSRHFNIHSWEKKYPCRYCEK
SEQ ID NO:313 YQCLACGKSFINYQFMSSHISVHSQDPSGDSKLYRLHPCRSIQLIRQYAY

PN7771

SEQ ID NO:314 LSLKENSRLLENELLENAEKL
SEQ ID NO:315 AEYENLTNKLQRNLENVLAE
SEQ ID NO:316 KFGDLDPSAEEFLQEERLT
SEQ ID NO:317 QMRNEYERQCRLQDQVDEL
SEQ ID NO:318 QSELEEVRAQGRVLRLPLKN
SEQ ID NO:319 QVVREKEKLRSDLDKAEKLKSLMASEVDDHAAIERNEYNLRKLDGEYK
SEQ ID NO:320 SEECPNLNMSIEAELVIEQMKEQHHRDICCLRLELEDKVRHYEKQLDETV

TSP3

SEQ ID NO:321 ILVELRDDIRDQVKEMSLIR
SEQ ID NO:322 NTIMECQVCGFHEQRSHCSP
SEQ ID NO:323 GPCPPGLQGNNGTHCSDINEC
SEQ ID NO:324 AHADPCFPGSSCINTMPGFH
SEQ ID NO:325 CEACPRTGYKGTQVSGVGIDY
SEQ ID NO:326 AVTNALHSILGEQTAKALVTQLTLFNQILVELRDDIRDQVKEMSLIRNTIM
SEQ ID NO:327 INECAHADPCFPGSSCINTMPGFHCEACPRTGYKGTQVSGVGIDYARASKQ

MDH1

SEQ ID NO:328 MSEPIRVLVTGAAGQIAYSL
SEQ ID NO:329 AFKDLDVAILVGSMPREGM
SEQ ID NO:330 VIIWGNHSSTQYPDVNHAKV
SEQ ID NO:331 VIIWGNHSSTQYPDVNHAKV
SEQ ID NO:332 SSAMSAAKAICDHVRDIWFG
SEQ ID NO:333 KKSVKVIVVGNPANTNCLTASKSAPSIPKENFSCLTRLDHNRAKAQIALK
SEQ ID NO:334 QYPDVNHAKVKLQGKEVGVYEALKDDSWLKGEFVTTVQQRGAAVIKARKL

GA17

SEQ ID NO:335 ESVVNSVSVSLLLILEPDKQE
SEQ ID NO:336 ALIESLCEKLVKFREGERPS
SEQ ID NO:337 LTIFVSAKLASYVKFYQNNK
SEQ ID NO:338 DFIDSLGLLHEQNMAKMRLL
SEQ ID NO:339 LQIGADDVEAFVIDAVRTKM
SEQ ID NO:340
MSVPAFIDISEEDQAAELRAYLKSKGAEISEENSEGLHVVDLAQIEACD
SEQ ID NO:341
LQIGADDVEAFVIDAVRTKMVYCKIDQTQRKVSHSTHRTFGKQRWQQL

CAPN4

SEQ ID NO:342 RQFRRLFQAQLAGDDMEVSAT
SEQ ID NO:343 MNILNKVVTRHPDLKTDGFG
SEQ ID NO:344 TCRSMVAVMDSDTTGKLGFE
SEQ ID NO:345 FDTDRSGTICSSLEPGAFEA
SEQ ID NO:346 AGFHLNEHLYNMIIIRRYSDE
SEQ ID NO:347
EAAGFHLNEHLYNMIIIRRYSDESGNMDFDNFISCLVRLDAMFRAFKSLDK
SEQ ID NO:348
MNILNKVVTRHPDLKTDGFGIDTCRSMVAVMDSDTTGKLGFEFKYLWNN

BAT3

SEQ ID NO:349 ERAPAQNPELTPGPAPAGPT
SEQ ID NO:350 PAPETNAPNHPSPAELYEV
SEQ ID NO:351 QELQRLESRLQPFLQRYYEV
SEQ ID NO:352 ESSAEGAPPPGPAPPATSH
SEQ ID NO:353 PRVIRISHQSVEPVMMHMN
SEQ ID NO:354
ERAPAQNPELTPGPAPAGPTPAPETNAPNHPSPAELYEV
SEQ ID NO:355
LGAAATTDDYNHNHEGREEDQRLINLVGESLRLLGNTFVALSDLRCNLACT

Of MSK-1 Interactors

ABLIM1 (778)

SEQ ID NO:356 GKTYHPNCFACTICKRPFPP
SEQ ID NO:357 GDRVTFNGRDCLCQLCAQPM
SEQ ID NO:358 SSSPKETTFSSNCAGCGRDI
SEQ ID NO:359 DIKNGQALLALDKQWHLGCF
SEQ ID NO:360
GKTYHPNCFACTICKRPFPPGDRVTFNGRDCLCQLCAQPMSSSPKETTF
SEQ ID NO:361
RDCLCQLCAQPMSSSPKETTFSSNCAGCGRDIKNGQALLALDKQWHLGCF

NICE-4

SEQ ID NO:362 TTQHSSTLSTQQNTLSSSTS
SEQ ID NO:363 HSSSSTFSTTSSTVSAPPPV
SEQ ID NO:364 VSVSSSLNSGSSLGLSLGSN
SEQ ID NO:365 NLPPGVPLLNPYIMAPGL
SEQ ID NO:366 LHAYPPQVGYDDLQMLQTR
SEQ ID NO:367
TSVESEANLHSSSSTFSTTSSTVSAPPVSVSSLNSGSSLGLSLGSNS
SEQ ID NO:368
GLLHAYPPQVGYDDLQMLQTRFPLDYYSIPFPPTPTPLTGRDGSLASNP

Of p38 ALPHA or MAPK14 (360) Interactors

CYT4

SEQ ID NO:369 PEPAEELSSGETEELQRIKWH
SEQ ID NO:370 KEKELCIGRKKFNMDPAKGI
SEQ ID NO:371 INLQVLQAFVDCHEFANLNL
SEQ ID NO:372 STDTCYVLSFSIIMLNTSLH
SEQ ID NO:373 DRMMEAFA TRYCLCNPGVFQ
SEQ ID NO:374
KFNMDPAKGIQYFIEHKLLTPDVQDIARFLYKGEGLNKTAIGTYLGERDP
SEQ ID NO:375
VDCHEFANLNQALRQFLWSFRLPGEAQKIDRMMEAFA TRYCLCNPGVF

JNK3A2

SEQ ID NO:376 RPKYAGLTFPKLFPDSLFP
SEQ ID NO:377 DSEHNKLKASQARDLLSKML
SEQ ID NO:378 VNSSES LPPSSSVNDI SSMS
SEQ ID NO:379 TDQTLASDTDSSLEASAGPL
SEQ ID NO:380 RNYVENRPKYAGLTFPKLFP
SEQ ID NO:381
VWYDPAEVEAPPPQIYDKQLDEREHTIEEWKELIYKEVMNSEEKTNGVV
SEQ ID NO:382
KLFPDSLFPADSEHNKLKASQARDLLSKMLVIDPAKRISVDDALQHPYIN

C-NAP1

SEQ ID NO:383 AAVVEARAQASAAGILEEDL
SEQ ID NO:384 RTARSALKKNEEVESERER
SEQ ID NO:385 AQALQEQQGELKVAQGKALQE
SEQ ID NO:386 LELEENHHKMECQQKLIKEL
SEQ ID NO:387 AQGKALQENLALLTQTLAER
SEQ ID NO:388
QKLTQREQIRELEKDRETQRNVLEHQLLELEKKDQMIESQRGQVQDLKK
SEQ ID NO:389
RGQIQELEKQREMQKAALLELLSDLKKRNQEVDLQQEQIQELEKCRSVLE

VCL (1066)

SEQ ID NO: 390 VTRLAKEVAKQCTDKRIRTN
SEQ ID NO: 391 LLQVCERIPTISTQLKILST
SEQ ID NO: 392 VKATMLGRTNISDEESEQAT
SEQ ID NO: 393 EMLVHNAQNLMQSVKETVRE
SEQ ID NO: 394 KIRTDAGFTLRWVRKTPWYQ
SEQ ID NO: 395
MSRLVGGSGTKRALIQCAKDIAKASDEVTRLAKEVAKQCTDKRIRTNLL
SEQ ID NO: 396
EQATEMLVHNAQNLMQSVKETVREAEAASIIRTDAGFTLRWVRKTPWYQ

PSF

SEQ ID NO: 397 VGNLPADITEDEFKRLFAY
SEQ ID NO: 398 KPAARKAFERCSEGVFLLTT
SEQ ID NO: 399 AQHGTFEYEYESQRWKSILDEM
SEQ ID NO: 400 EKQQREQVEKNMKDAKDKE
SEQ ID NO: 401 LSLLRRPGEKTYTQRCRLFG
SEQ ID NO: 402
QLRVRFATHAALSVRNLSPYVSNELLEEAFSQFGPIERAVVIVDDRGRS
SEQ ID NO: 403
KPAARKAFERCSEGVFLLTTPRPVIVEPLEQLDDEDGLPEKLAQKNPMY

Of AKT1 Interactors

FNTA

SEQ ID NO: 404 IADILNQDAKNYHAWQHRQW
SEQ ID NO: 405 VIQEFLWDNELQYVDQLLK
SEQ ID NO: 406 YFVISNTTGYNDRAVLEREV
SEQ ID NO: 407 QYTLEMIKLVPHNESAUNYL
SEQ ID NO: 408 KGILQDRGLSKYPNLLNQLL
SEQ ID NO: 409
YLKGILQDRGLSKYPNLLNQLLDLQPSHSSPYLIAFLVDIYEDMLENQCD
SEQ ID NO: 410
DILNQDAKNYHAWQHRQWVIQEFLWDNELQYVDQLLKEDVRNNNSVWNQR

TPRD

SEQ ID NO: 411 NRNSDSAGPFAVPDHLRQDV
SEQ ID NO: 412 EALYDQHSNEYVVRNKKLWD
SEQ ID NO: 413 MNPKQKCSTLYDYFSQFLEE
SEQ ID NO: 414 HGPLDMSNKMFSAEYEFFPE
SEQ ID NO: 415 ETRQILEKAGGLKPFLLGCP
SEQ ID NO: 416
EETRQILEKAGGLKPFLLGCPRFVVIDNCIALKKVASRLKKRKKNIKT
SEQ ID NO: 417
NRNSDSAGPFAVPDHLRQDVEEFEALYDQHSNEYVVRNKKLWDMNPQKC

PPL

SEQ ID NO:418 EFHNSKSSKELDFLREENHK
SEQ ID NO:419 DLKRLSKDKDLEIDELQKRL
SEQ ID NO:420 DWNMFVKLRSQECDWEEISV
SEQ ID NO:421 RYVNKDMSIQELAVLVSGQK
SEQ ID NO:422 VAVKREQRENHLRRSIVVIH
SEQ ID NO:423
LREENHKLQLERQNLQLETRRLQSEINMAATETRDLRNMTVADSGTNHDS
SEQ ID NO:424
RSQECDWEEISVKGPNGESSVIHDRKSGKKFSIEEALQSGRLTPAQYDRY

GOLGIN-84

SEQ ID NO:425 STEKNSLVFQLERLEQQMNS
SEQ ID NO:426 DTETNLAGMYGKVRKAASSI
SEQ ID NO:427 YPIARVFVIIYMAILHLWVM
SEQ ID NO:428 VMIVLLTYTPEMHHDQPYGK
SEQ ID NO:429 MSGIDNGEGTRLRNVPVLFN
SEQ ID NO:430
STEKNSLVFQLERLEQQMNSASGSSSSNGSSINMSGIDNGEGTRLRNVPVL
SEQ ID NO:431
DTETNLAGMYGKVRKAASSIDQFSIRLGIFLRRYPIARVFVIIYMAILHL

of AKT2p55 (481) Interactors

CLIC1

SEQ ID NO:432 RYPKLAALNPESNTAGLDIF
SEQ ID NO:433 AKFSAYIKNSNPALNDNLEK
SEQ ID NO:434 VDETSAADEGVSKFLDGN
SEQ ID NO:435 ELTLADCNLLPKLHIVQVVC
SEQ ID NO:436 NAYAREEFASTCPDDEEIEL
SEQ ID NO:437
ESNTAGLDIFAKFSAYIKNSNPALNDNLEKGLLKALKVLDNYLTSPLEE
SEQ ID NO:438
NLLPKLHIVQVVCKKYRGFTIPEAFRGVHRYLSNAYAREEFASTCPDDEE

AKR7A2

SEQ ID NO:439 GLGLGGGDCRVKIATKANPW
SEQ ID NO:440 EETLHACQRLHQEGKFVELG
SEQ ID NO:441 ELFPCLRHFGLRFYAYNPLA
SEQ ID NO:442 FEAIALVEKALQAAYGASAP
SEQ ID NO:443 TEEGPLEPAVVDAFNQAWHL
SEQ ID NO:444
HQEGKFVELGLSNYASWEVAEICTLCKSNGWILPTVYQGMYNATTRQVET
SEQ ID NO:445
LQAAYGASAPSVTSAALRWMYHHSQGAHDAVILGMSSLEQLEQNLAA

TPRD

SEQ ID NO:446 SNRNSDSAGPFAVPDHLRQD
SEQ ID NO:447 EALYDQHSNEYVVRNKKLWD
SEQ ID NO:448 MNPQKQKCSTLYDYFSQFLEE
SEQ ID NO:449 HGPLDMSNKMFSAEYEFFPE
SEQ ID NO:450 ETRQILEKAGGLKPFLLGCP
SEQ ID NO:451
EETRQILEKAGGLKPFLLGCPRFVVIDNCIALKKVASRLKKRKKNIKT
SEQ ID NO:452
FAVPDHLRQDVVEFALYDQHSNEYVVRNKLWDMNPQKCSTLYDYFSQF

Of P90RSK or RPS6KA1 Interactors

UNR (798)

SEQ ID NO:453 GQSPTGSVCYERNGEVFYLT
SEQ ID NO:454 KKQARCQGVVCAMKEAFGFI
SEQ ID NO:455 VATDVRLLPQGTIVFEDISI
SEQ ID NO:456 DKDTKSKVTLLEGDHVRFNI
SEQ ID NO:457 HSHSDHRFLGTVEKEATFSN
SEQ ID NO:458
GFIKCVDRDVRMFFHFSEILDGNQLHIADEVEFTVVPDMLSAQRNHAIRI
SEQ ID NO:459
CAMKEAFGFIERGDVVKEIFFHYSEFKGDLETQPGDDVEFTIKDRNGKE

Of IKK-alpha Interactor

GBDR1

SEQ ID NO:460 HKLIHAASERVLSARTILE
SEQ ID NO:461 ENIQDQDVLLLIKRAAPSPL
SEQ ID NO:462 PKMADVSAEKKQDQKAPD
SEQ ID NO:463 KEAILRATANLPSYNMDRAA
SEQ ID NO:464 VQTNMRDFQTELRKILVSLI
SEQ ID NO:465
KSITHHKLIHAASERVLSARTILEENIQDQDVLLLIKRAAPSPLPKMAD
SEQ ID NO:466
MDRAAVQTNMRDFQTELRKILVSLIEVAQKLLALNPDAVELFKKANAMLD

Of IKK-beta Interactors

KIAA0614

SEQ ID NO:467 SSQESLDISLCSTGSLGSLG
SEQ ID NO:468 VVEIRSRAKIEKIRASFNN
SEQ ID NO:469 GLEDYFNDKSIKGEKLVPGA
SEQ ID NO:470 QTPGNLLHLFFTNRPPKKV

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SEQ ID NO:471 PKFDKSKYSKAGKEQHPVKV
SEQ ID NO:472
EKIRASLFNNNDLIGLSSLGEDELMEMSTEEILTVSVNQSLFDTQGSP
SEQ ID NO:473
SIKGEKLVPGAREVLTEIFKSCAHSEQTLSLTPAKPIRVSDIYLSKEQIN

LDHM

SEQ ID NO:474 TESRSSFSQHARTSGRVAVE
SEQ ID NO:475 YRFPPKFTLKAGQVVTIWAA
SEQ ID NO:476 RKLVRSTVVEDDEDEDGDD
SEQ ID NO:477 RLRSPGGRTHLLWLFCLQCH
SEQ ID NO:478 LRGQSGHPLLPGQLQPPNP
SEQ ID NO:479
RSSFSQHARTSGRVAVEEVDEEGKFVRLRNKSNEQSMGNWQIKRQNGDD
SEQ ID NO:480
HLLWLFCLQCHGHSQLPQCGGQWGWLQRGQSGHPLLPGQLQPPNPEPPE

EIF3S10

SEQ ID NO:481 PDFIMAKQVEQLEKEKKELQ
SEQ ID NO:482 DMDLWEQQEERITTMQLER
SEQ ID NO:483 ERLAERHNRLEERKRQRKE
SEQ ID NO:484 ERRITYREKEEEEQRRAEE
SEQ ID NO:485 QMLKEREERERAERAKREEE
SEQ ID NO:486
RITYREKEEEEQRRAEEQMLKEREERERAERAKREEELREYQERVKKLE
SEQ ID NO:487
MAKQVEQLEKEKKELQERLKNQEKKIDYFERAKLEEIPLIK SAYEEQRI

SLAP-2

SEQ ID NO:488 QADNDFTNERLTALQENQTR
SEQ ID NO:489 MDEQDLNEPLAKVSLLKDDL
SEQ ID NO:490 EEERKAYRNQVEESTKQIQV
SEQ ID NO:491 LHQAAAKVASERDTDIASLQ
SEQ ID NO:492 EDQQREEATRLQGELEKLRK
SEQ ID NO:493
LAKVSLLKDDLQGAQSEIEAKQEIQHLRKELIEAQELARTSKQKCFELQA
SEQ ID NO:494
LHQAAAKVASERDTDIASLQEELKKVRAELERWRKAASEYEKEITSQNS

SART-1

SEQ ID NO:495 LYSARDLQGLTVEHAIDSFR
SEQ ID NO:496 EKNVELRKKKPDYLPYAED
SEQ ID NO:497 SVDDLAQQKPRSILSKYDEE
SEQ ID NO:498 LEGERPHSFRLEQGGTADGL
SEQ ID NO:499 RERELEEIRAKLRLQAQSL
SEQ ID NO:500
LYSARDLQGLTVEHAIDSFREGETMILTLKGVLQEEEDVLVNVNLVDK

SEQ ID NO:501
EIRAKLRLQAQSLSTVGPRYLASEYLTPEEMVTFKTKRRVKKIRKKEKEV

GBDR1

SEQ ID NO:502 HKLIHAASERVLSARTILE
SEQ ID NO:503 ENIQQDQDVLLLIKRAAPSPL
SEQ ID NO:504 PKMADVSAEKKQDQKAPD
SEQ ID NO:505 KEAILRATANLPSYNMDRAA
SEQ ID NO:506 VQTNMRDFQTELRKILVSLI
SEQ ID NO:507
ITHHKLIHAASERVLSARTILEENIQDQDVLLLIKRAAPSPLPKMADVS
SEQ ID NO:508
APDKEAILRATANLPSYNMDRAAVQTNMRDFQTELRKILVSLIEVAQKLL

Of IKK-gamma Interactors

ITRAF

SEQ ID NO:509 TENYEQRIREQQEQLSLQQT
SEQ ID NO:510 ETSARSLGSPLLHERGNIEK
SEQ ID NO:511 NDSTFLHSTPERPGILSPAT
SEQ ID NO:512 AIRGPQQPIWKPFPNQDSDS
SEQ ID NO:513 VVLSGTDSELHIPRVCEFCQ
SEQ ID NO:514
QLLLVNSTQDNNYGCVPLLESDTRKNTLTLAQPKVISGIAREKLPKV
SEQ ID NO:515
QNLKTTDKTKPSNLVNTCIRTLDRACLPPGDHNALYVNSFPPLDPSDA

Of IKK-i Interactors

ITRAF

SEQ ID NO:516 QLLLVNSTQDNNYGCVPLLE
SEQ ID NO:517 ETSARSLGSPLLHERGNIEK
SEQ ID NO:518 IQCTDKTDKQEALFTPQAKD
SEQ ID NO:519 NDSTFLHSTPERPGILSPAT
SEQ ID NO:520 QNLKTTDKTKPSNLVNTCIR
SEQ ID NO:521
AIRGPQQPIWKPFPNQDSDSVVLSGTDSELHIPRVCEFCQAVFPPSITSR
SEQ ID NO:522
QDNNYGCVPLLESDTRKNTLTLAQPKVISGIAREKLPKVRRQEVSPP

NUMA1

SEQ ID NO:523 STQAALQAMEREAEQMGNEL
SEQ ID NO:524 ALEKAARAELEMRLQNALNE
SEQ ID NO:525 QRVEFATLQEALAHALTEKE
SEQ ID NO:526 GKDQELAKLRGLEAAQIKEL

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SEQ ID NO:527 EAAQIKELEELRQTVKQLKE
SEQ ID NO:528
AMEREAEQMGNELERLRAALMESQGQQQEERGQQEREVARLTQERGRAQA
SEQ ID NO:529
VEFATLQEALAHALTEKEGKDQELAKLRGLEAAQIKELEELRQTVKQLKE

SPA1

SEQ ID NO:530 LEDEWQAISEIASTCNTILE
SEQ ID NO:531 GTPKSDAEPEPGNLSEKVSH
SEQ ID NO:532 LESMLRKLQEDLQKEKADRA
SEQ ID NO:533 ALEEEVRSLRHNNRRLQAES
SEQ ID NO:534 ESAATRLLLASKQLGSPTAD
SEQ ID NO:535
LEDEWQAISEIASTCNTILESLSREGQPIPEGDPKGTPKSDAEPEPGNL
SEQ ID NO:536
QKEKADRAALEEEVRSLRHNNRRLQAESESAATRLLLASKQLGSPTADLA

FYCO1

SEQ ID NO:537 SLVSSYLQTQEMVSNFDLNS
SEQ ID NO:538 QLEVREKQLQERMQQLDREN
SEQ ID NO:539 QWEVTQATQNTVKELQTCLQ
SEQ ID NO:540 KNQHLASFPGLAMAQQKAD
SEQ ID NO:541 NRQQSAQLEQLVKELQLKED
SEQ ID NO:542
LKEDARASLERLVKEMAPLQEELSGKGQEADQLWRRLQELLAHTSSWEET
SEQ ID NO:543
AVSQQGEQLQTERERGRRTAAEDNVRLTCLVAELQKQWEVTQATQNTVKEL

of Survivin/BIRK5 Interactors

DNCL1

SEQ ID NO:544 MCDRKAVIKNADMSEEMQQD
SEQ ID NO:545 SVECATQALEKYNIEKDIAA
SEQ ID NO:546 HIKKEFDKKYNPTWHCIVGR
SEQ ID NO:547 NFGSYVTTHEKHFIFYFYLGQ
SEQ ID NO:548
AHIKKEFDKKYNPTWHCIVGRNFGSYVTTHEKHFIFYFYLGQVAILLFKSG
SEQ ID NO:549
MCDRKAVIKNADMSEEMQQDSVECATQALEKYNIEKDIAAHIKKEFDKKY

ACTB

SEQ ID NO:550 KRGILTLKYPIEHGIVTNWD
SEQ ID NO:551 QIMFETFNTPAMYVAIQAVL
SEQ ID NO:552 GRDLTDYLMKILTERGYSFT
SEQ ID NO:553 AGRDLTDYLMKILTERGYSF
SEQ ID NO:554 GGTTMYPGIADRMQKEITAL

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SEQ ID NO:555

IEHGIVTNWDDMEKIWHHTFYNELRVAPEEHPLLTEAPLNPKANREKMT

SEQ ID NO:556

KIKIIAPPERKYSWIGGSILASLSTFQQMWISKQEYDESGPSIVHRKCF

KU70

SEQ ID NO:557 FQDMMGHGSDYSLSEVLWVC

SEQ ID NO:558 AKASRARTKAGDLRDTGIFL

SEQ ID NO:559 DLLRKVRAKETRKRALSRLK

SEQ ID NO:560 RTFNTSTGLLLLPSDTKRSQ

SEQ ID NO:561 YLRPSLFVYPEESLVIGSST

SEQ ID NO:562

DLMHLKKPGGFDISFYRDIISIAEDEDLRVHFEESSKLEDLLRKVRAKE

SEQ ID NO:563

ISVGIYNLVQALKPPPPIKLYRETNEPVKTTRTFNTSTGLLLLPSDTKR

COPP

SEQ ID NO:564 NQKAAESLADPTEYENLFPG

SEQ ID NO:565 LKEAFVVEEWVKETHADLWP

SEQ ID NO:566 AKQYPLVTPNEERNVMEEGK

SEQ ID NO:567 DFQPSRSTAQQELDGKPASP

SEQ ID NO:568 TPVIVASHTANKEEKSLEL

SEQ ID NO:569

VVKLWRENLSKVNQKAAESLADPTEYENLFPGGLKEAFVVEEWVKETHADL

SEQ ID NO:570

PASPTPVIVASHTANKEEKSLELEVLDLNLELEDIDTTDINLDEDILDD

OSTP

SEQ ID NO:571 MRIAVICFCLLGITCAIPVK

SEQ ID NO:572 QADSGSSEEKQLYNKYPDAV

SEQ ID NO:573 PDAVATWLNPDPSQKQNLLA

SEQ ID NO:574

MRIAVICFCLLGITCAIPVKQADSGSSEEKQLYNKYPDAVATWLNPDPSQ

SLC8A1

SEQ ID NO:575 NFLDGALVLEVDERDQDDE

SEQ ID NO:576 RAFYRIQATRLMTGAGNILK

SEQ ID NO:577 NCGTVALTIIRRGGDLTNTV

SEQ ID NO:578 DDDIFEEDENFLVHLSNVKV

SEQ ID NO:579 TFEPPVTHVSESIGIMEVKV

SEQ ID NO:580

RREMARILKELKQHPDKEIEQLIELANYQVLSQQQKSRAFYRIQATRLM

SEQ ID NO:581

TATVTIFDDDHAGIFTFEPPVTHVSESIGIMEVKVLRSGARGNVIVPYK

A2-CAT

SEQ ID NO:582 AASVEQATQNFLEKGEQIAK

SEQ ID NO: 583 KKMVKLNVAARRQQELKDP
SEQ ID NO: 584 PSLEERLESIISGAALMADS
SEQ ID NO: 585 FREHANKLVEVANLACSI
SEQ ID NO: 586 SLCPQVINAALTTLAARPQSK
SEQ ID NO: 587
GTMVRAARALLSAVTRLLILADMADVMRLLSHLKIVEEALEAVKNATNEQ
SEQ ID NO: 588
PLNIAIDKMTKKTRDLRRQLRKAVMDHISDSFLETNVPLLVLIEAAKSGN

Of COX1 Interactors

THRSP

SEQ ID NO: 589 PSLLRDVQLSGPGGQAQAEA
SEQ ID NO: 590 TMLKAICVDVDHGLLPREEW
SEQ ID NO: 591 QAKVAGSEENGTAETEEVED
SEQ ID NO: 592 ESASGELDLEAQFHLHFSSL
SEQ ID NO: 593 HHILMHLTEKAQEVT
SEQ ID NO: 594
PSLLRDVQLSGPGGQAQAEA
SEQ ID NO: 595 ENGTAETEEVED
ESASGELDLEAQFHLHFSSL
HHILMHLTEKAQEVT
RYQE

OPA1

SEQ ID NO: 596 LVKLAPDFDKIVESLSLLKD
SEQ ID NO: 597 DFFTSGSPEETAFRATDRGS
SEQ ID NO: 598 KELRKLVQKDDKG
SEQ ID NO: 599 IHHRKL
SEQ ID NO: 600 KTSVLEMIAQARIFPRGS
SEQ ID NO: 601 GCTVSPE
SEQ ID NO: 602 HEIELRMRKNV
SEQ ID NO: 603 FRKVSDKEKIDQLQEELLHTQLKYQRILERLEKEN
SEQ ID NO: 604 KELRKLVQKDDKG
SEQ ID NO: 605 IHHRKL
SEQ ID NO: 606 KTSVLEMIAQARIFPRGS
SEQ ID NO: 607 GCTVSPE
SEQ ID NO: 608 HEIELRMRKNV
MTRSPVKVTLSEGPHHVALFKDSSREFDLTKEEDLAALRHEIELRMRKNV

Of SET(277) Interactor

PN12218

SEQ ID NO: 603 TKGHSYPPGPAEVSRRLRRCR
SEQ ID NO: 604 SAQPSPERQPSPLEPRPVSP
SEQ ID NO: 605 AALDALDKAQRQLQACKRRE
SEQ ID NO: 606 AREKRAQADARQTLKEHVQD
SEQ ID NO: 607 EKRAQADARQTLKEHVQDFA
SEQ ID NO: 608 PLGAQADEAGCSAQPSPERQPSPLEPRPVSP
SEQ ID NO: 609 SAYMLRLPPPAGAYIQNEH
PLGAQADEAGCSAQPSPERQPSPLEPRPVSP
SEQ ID NO: 610 SAYMLRLPPPAGAYIQNEH

Of ZFP36 Interactors

CIN85

SEQ ID NO:610 AIVEFDYQAQHDDELTISVG
SEQ ID NO:611 KKDPLTNKAPEKPLHEVPSG
SEQ ID NO:612 VGDIIIEVVGEVEEGWWEGLV
SEQ ID NO:613 SESDGGDSSTKSEGANGTV
SEQ ID NO:614 EKTIGKKLPATTATPDSSKT
SEQ ID NO:615
HDDELTISVGEIITNIRKEDGGWWEGQINGRRGLFPDNFVREIKKEMKKD
SEQ ID NO:616
TKSEGANGTVATAAIQPKKVKGVGFGLIFKDKPIKLPRSIIEVENDFLPV

PN13734

SEQ ID NO:617 EHGEPEFIQACMTFLKRRCP SILGG
SEQ ID NO:618 QACAGSVSQELSETILTMVANCSNVMNKA
SEQ ID NO:619 GMTSLSIGGSAAPHTQSMQGFPPNLGS
SEQ ID NO:620 LPVGGLGTGSILTGIGT GALGLPAVNNDPF
SEQ ID NO:621 ANQHFSKEIDDEANSYFQRIYNHP
SEQ ID NO:622
FEEYRFFFQYPDKELHITACLFGGIEKGLVTYMALGLALRYVLEALRKP
SEQ ID NO:623
DTLLVATDQTERIVEPPENIQEKIAFIFNNLSQSNTQKVEELKETVKEE

Of TIAL1(375) Interactors

FUBP1

SEQ ID NO:624 MADYSTVPPPSSGSAGGGGGGGGGGVN
SEQ ID NO:625 EQISRIQQESGCKIQIAPDS
SEQ ID NO:626 EQISRIQQESGCKIQIAPDS
SEQ ID NO:627 PYKVQQAKEMVLELIRDQGG
SEQ ID NO:628 RGRGQGNWNMGPPGGLQEFN
SEQ ID NO:629
MGPYNPAPYNPGPPGPAPHGPPAPYAPQGWGNAYPHWQQAPPDAKAGT
SEQ ID NO:630
GGQKRPLEDGDQPDAKKVAAPQNDSFGTQLPPMHQQQSR SVMTEEYKVPDG

Of CLIC1 Interactors

LRP1

SEQ ID NO:631 EWLCLLSPSGPVCTCPNGKRLDN
SEQ ID NO:632 NLQCFNNGGSCFLNARRQPKCRCQPRYT
SEQ ID NO:633 FTGPKCTQQVCAGYCANNSTCTVNQ

SEQ ID NO: 634 AADGSRQCRCCTAYFEGSRCEVNKCSRLE
SEQ ID NO: 635 CSNGGSTMNSKMMPECQCPCPHMTGP
SEQ ID NO: 636 VFWYKRRVQGAKGFQHQRMNTNGAMNVEIGNPTYKMYEGGEPPDVGLLDA
SEQ ID NO: 637 CFLNARRQPCKRCQPRYTGDKELDQCWEHCRNGGTCAASPSGMPTCRCP

FUS (526)

SEQ ID NO: 638 MASNDYTQQATQSYGAYPTQ
SEQ ID NO: 639 PGQGYSQQSSQPYGQQSYSG
SEQ ID NO: 640 YSQSTDTSGYGQSSYSSYGQ
SEQ ID NO: 641 SQNTGYGTQSTPQGYGSTGG
SEQ ID NO: 642 YGSSQSSQSSYGQQSSYPGY
SEQ ID NO: 643 TGYGTQSTPQGYGSTGGYGSQSQQSSYGQQSSYPGYGQQPAPSSTGSY
SEQ ID NO: 644 MASNDYTQQATQSYGAYPTQPGQGYSQQSSQPYGQQSYSGYSQSTDTSY

FUS (525)

SEQ ID NO: 645 MASNDYTQQATQSYGAYPTQ
SEQ ID NO: 646 PGQGYSQQSSQPYGQQSYSG
SEQ ID NO: 647 YSQSTDTSGYGQSSYSSYGQ
SEQ ID NO: 648 SQNSYGTQSTPQGYGSTGGY
SEQ ID NO: 649 GSSQSSQSSYGQQSSYPGY
SEQ ID NO: 650 NSYGTQSTPQGYGSTGGYGSQSQQSSYGQQSSYPGYGQQPAPSSTGSY
SEQ ID NO: 651 ASNDYTQQATQSYGAYPTQPGQGYSQQSSQPYGQQSYSGYSQSTDTSY
G

of NCOA2 Interactors

XE169

SEQ ID NO: 652 DTKFLCPLCMRSRRPRLETI
SEQ ID NO: 653 LLVALQRLPVRLPEGEALQC
SEQ ID NO: 654 PEEPPNYPAAPASDPLREGS
SEQ ID NO: 655 LPQLTGPVLELPEATRAPLE
SEQ ID NO: 656 EELMMEGDLLEVTLDENHSI
SEQ ID NO: 657 VRLPEGEALQCLTERAISWQGRARQALASEDVTALLGRLAELRQLQAEP
SEQ ID NO: 658 DTKFLCPLCMRSRRPRLETI ALLVALQRLPVRLPEGEALQCLTERAISW

NAG4

SEQ ID NO: 659 QDYPYVMADSLLDVLTKGHH
SEQ ID NO: 660 MSLPEDEGHTRTLDTAKEME
SEQ ID NO: 661 IFQKKLDETTRLLRELQEAQ

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SEQ ID NO:662 PPNMICLLGPSYREMHLAEQ
SEQ ID NO:663 TPGDIVSTYGVRKAMGISIP
SEQ ID NO:664
MSLPEDEGHTRTLDTAKEMEITEVEPPGRLDSSTQDRLIALKAVTNFGVP
SEQ ID NO:665
IFQKKLDETTRLLRELQEAQNERLSTRPPPNNICLLGPSYREMHLAEQVT

ESRRA

SEQ ID NO:666 VAGGPRKTAAPVNALVSHLL
SEQ ID NO:667 VEPEKLYAMPDPAGPDGHL
SEQ ID NO:668 AVATLCDLFREIVVTISWA
SEQ ID NO:669 KSI PGFSSLSLSQMSVLQS
SEQ ID NO:670 VWMEVLVLGVAQRSLPLQDE
SEQ ID NO:671
VAGGPRKTAAPVNALVSHLLVVEPEKLYAMPDPAGPDGHLPAVATLCDLF
SEQ ID NO:672
IPGFSSLSLSQMSVLQSVWMEVLVLGVAQRSLPLQDELAF AEDLVLD EEE

CTNNB1

SEQ ID NO:673 LHYGLPVVVKLLHPPSHWPL
SEQ ID NO:674 VRAHQDTQRRTSMGGTQQQF
SEQ ID NO:675 QLLYSPNIQRVAAGVLCE
SEQ ID NO:676 IEAEGATAPL TELLHSRNEG
SEQ ID NO:677 GALHILARDVHN RIVIRGLN
SEQ ID NO:678
QRRTSMGGTQQQFVEGVRMEEIVEGCTGALHILARDVHN RIVIRGLNTIP
SEQ ID NO:679
CELAQDKEAAEAI EAEGATAPL TELLHSRNEG VATYAAVLFRMSEDKPQ

CIT

SEQ ID NO:680 SKIDWNNIRNS PPPFVPTLK
SEQ ID NO:681 YSKALGILGRSES VVSGLDS
SEQ ID NO:682 EVEAVLSQKEVELKASETQR
SEQ ID NO:683 DIREQSRKLQEIKEQEQYQAQ
SEQ ID NO:684 RKATECQHKLLKAKDQGKPE
SEQ ID NO:685
IRQAKERAERELEKLQNREDSSEGIRKKLVEAEERRHSLENKVKRLETME
SEQ ID NO:686
IQTKSQQIQQMADKILELEEKHREAQVSAQHLEVHLKQKEQHYEEKIKVL

HSBP1

SEQ ID NO:687 MSLFDLFRGFFGFPGRSHR
SEQ ID NO:688 FGF GFSFSPGGGIRFHDNFG
SEQ ID NO:689 EGQTLRDSMLKYPDSHQPRI
SEQ ID NO:690 TRENDLDSQVSQEGLGPVL
SEQ ID NO:691 TRHEADSSPRGDPE SPRPPA

SEQ ID NO:692
PKSYFKSISVTKITKPDGIVEERRTVVDSEGRTEVVTRHEADSSPRGDP
SEQ ID NO:693
EEGGSWRRGNPRFHSPQHPPEFGFGFSFSPGGGIRFHNDNFGFDDLVRDF

ROCK2

SEQ ID NO:694 SKQIQQLESNNRDLQDKNCL
SEQ ID NO:695 LQDKNCLLETAKLKLEKEFI
SEQ ID NO:696 LAKVELEKRQLQERFTDLEK
SEQ ID NO:697 IYESIEEAKSEAMKEMEKKL
SEQ ID NO:698 QKDVLNEDVRNLTLKIEQET
SEQ ID NO:699
ERRDRTHGSEIINDLQGRICGLEEDLKNGKILLAKVELEKRQLQERFTDL
SEQ ID NO:700
SQQKINELLKQKDVLNEDVRNLTLKIEQETQKRCLTQNDLKMQTQQVNTL

TILP (392)

SEQ ID NO:701 LKVLNEQLQSKEVELLQLRT
SEQ ID NO:702 KECSDLKIELQKAKQTDPYQ
SEQ ID NO:703 AELLRKLKTSTAIKKACAPV
SEQ ID NO:704 PLPGDVKVLSEKAILQSWT
SEQ ID NO:705 AFGETKTKTLPNLPPLHY
SEQ ID NO:706
ETQQVMRNLNPPSSNWEVEKLSCLKIHGLEQELEMRKECSDLKIELQK
SEQ ID NO:707
STKLHLMNFTATYTRHPPLLNGKALCHTTSSPLPGDVKVLSEKAILQSW

LRRFIP2a

SEQ ID NO:708 YMQGLKELKESLSEVEEKYK
SEQ ID NO:709 NEEKSKELERQKHMCSVLQH
SEQ ID NO:710 LWKDKKIGALEKQKEYIACL
SEQ ID NO:711 EPVAGAITVVSQEAAQVLES
SEQ ID NO:712 NDGTVGDLAGLQNGSDLQFI
SEQ ID NO:713
LKEGLRQRDELIEEKQRMQQKIDTMTKEVFDLQETLLWKDKKIGALEKQK
SEQ ID NO:714
SAGEGPLDVRRLRKLAGEKEELLSQIRKLKLQLEERQCSRNDGTVGDLA

PSAP

SEQ ID NO:715 SLQKH LAELNHQKQLESNKI
SEQ ID NO:716 PQDGP RSKPQPKDNGDVCQD
SEQ ID NO:717 CKNYI SQYSEIAIQMMMHMQ
SEQ ID NO:718 PIKKHEVPAKSDVYCEVCEF
SEQ ID NO:719 NKTEKEILD AFDKMCSKLPK
SEQ ID NO:720
MVTDIQTA VRTNSTFVQALVEHVKEEC DR LGPGMADICK NYI SQYSEIAI

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SEQ ID NO:721

QKQLESNKIPELDMTEVVAPFMANIPLLLYPQDGPRSKPQPKDNGDVCQD

Of ER-beta and ESR1 Interactor

NOTCH2

SEQ ID NO:722 MPALRPALLWALLALWLCCA

SEQ ID NO:723 YCQHRDPCEKNRCQNGGTCV

SEQ ID NO:724 CHMLSRDTYECTCQVGFTGK

SEQ ID NO:725 TDVNECDIPGHCQHGGTCLN

SEQ ID NO:726 VNGGTCRQTGDFTFECNCLP

SEQ ID NO:727

TPAHALQCRDGYEPCVNEGMCVTYHNGTGYCKCPEGFLGEYCQHRDPCEK

SEQ ID NO:728

SGFTGEDCQYSTSHPCFVSRPCLNNGTCHMLSRDTYECTCQVGFTGKECQ